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PATENT

## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant:

Heng Tsai, et al.

Serial No:

10/583,179

Filed:

June 16, 2006

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Title:

PROTEIN SEPARATION DEVICE

Examiner:

Group Art Unit: 2161

Docket No:

490352-3004

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2. After the period specified in paragraph (1) hereinabove of this section, but is being filed before the mailing date of either a final action under 37 CFR 1.113, or a notice of

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Respectfully submitted,

Bv

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Sheet 1 of 4

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Complete if Known					
Application Number	PCT/SG2004/000417				
Filing Date	December 17, 2004				
First Named Inventor	Heng Hang TSAI et al.				
Art Unit	not assigned yet				
Examiner Name	not assigned yet				
Attorney Docket Number	490352-3004/US				

U.S. PATENT DOCUMENTS								
Examiner	Cite	Document Number	Publication Date	Name of Patentee or	Page, Columns, Lines, Where			
Initials *	No. ¹	Number-Kind code 2 (if known)	MM-DD-YYY	Applicant of Cited Document	Relevant Passages or Relevant Figures Appear			
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	FOREIGN PATENT DOCUMENTS								
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	B1	WO 2002/083725, A1	10-24-2002	RNA INC. et al.					
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December 17, 2004

Hen Hang TSAI et al.

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Filing Date

Art Unit

First Named Inventor

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Sheet	2	of	4	Attorney Docket Number	490352-3004/US

		NON PATENT LITERATURE DOCUMENTS	
Examiner Initials *	Cite No. 1	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, pages), volume-issue number(s), publisher, city and /or country where published.	T <sup>2</sup>
	C1	AGARD, D.A., "To fold or not to fold", Science, (1993) p. 1903-1904, Vol. 260.	
	C2	ALTSCHUL, S.F., et al., "Basic Local Alignment Search Tool", J. Mol. Biol., (1990) p. 403-410, Vol. 215.	
	C3	AUSUBEL, et al. (eds), Current Protocols in Molecular Biology, (1989) p. 2.10.3, Vol. 1, Green Publishing Associates, Inc. and John Wiley & Sons, Inc., New York.	
	C4	BHUTANI, N., et al., "Chaperonins as protein-folding machines", Current Science, (2002) p. 1337-1351, Vol. 83, No. 11.	
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	C11	GÉMINARD, et al., "Characteristics of the interaction between Hsc70 and the transferrin receptor in exosomes released during reticulocyte maturation", J. Biological Chem., (2001) p. 9910-9916, Vol. 276.	
	C12	HARTL, F.U., et al., "Molecular Chaperones in the Cytosol: from Nascent Chain to Folded Protein", Science, (2002) p.1852-1858, Vol. 295.	
	C13	HEMMINGSEN, S.M., "What is a chaperonin?", Nature, (1992) p. 650, Vol. 357, No. 6380.	
	C14	HOURY, W.A., "Mechanism of substrate recognition by the chaperonin GroEL", Biochem. Cell Biol., (2001) p. 569-577, Vol. 79, © 2001 NRC Canada.	

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s	Substitute for form	1449/PTO		Complete if Known		
				Application Number	PCT/SG2004/000417	
n		MONTA	DISCLOSURE	Filing Date	December 17, 2004	
				First Named Inventor	Hen Hang TSAI et al.	
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(Use as many sheets as necessary)				Examiner Name	not yet assigned	
Sheet	3	of	4	Attorney Docket Number	490352-3004/US	

		NON PATENT LITERATURE DOCUMENTS	
Examiner Initials *	Cite No. <sup>1</sup>	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, pages), volume-issue number(s), publisher, city and /or country where published.	T <sup>2</sup>
	C15	HYNES, et al., "Individual subunits of the eukaryotic cytosolic chaperonin mediate interactions with binding sites located on subdomains of β-actin", J. Biological Chem., (2000) p. 18965-18994, Vol. 275.	
	C16	KINBARA, K., et al., "RAS GTPases: INTEGRINS' FRIENDS OR FOES?", Nature Reviews: Molecular Cell Biology, (2003) p. 767-776, Vol. 4.	
·	C17	KLUMPP, M., et al., "Structure of the Substrate Binding Domain of the Thermosome, an Archaeal Group II Chaperonin", Cell, (1997) p. 263-270, Vol. 91.	
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	C20	NEEDLEMAN, et al., "A General Method Applicable to the Search for Similarities in the Amino Acid Sequence of Two Proteins", J. Mol. Biol., (1970) p. 443-453, Vol. 48.	
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	C22	PEARSON, W.R., et al., "Improved tools for biological sequence comparison", Proc. Natl. Acad. Sci. USA, (1988) p. 2444-2448, Vol. 85.	
	C23	SAKIKAWA, C., et al., "On the Maximum Size of Proteins to Stay and Fold in the Cavity of GroEL underneath GroES", J. Biol. Chem., (1999) p. 21251-21256, Vol. 274, No. 30, Issue of July 23.	
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	C25	SMITH, et al., "Comparison of Biosequences", Adv. Appl. Math., (1981) p. 482-489, Vol. 2.	
	C26	STANIFORD, R.A., et al., "The stability and hydrophobicity of cytosolic and mitochondrial malate dehydrogenases and their relation to chaperonin-assisted folding", FEBS Letters, (1994) p. 129-135, Vol. 344.	
	C27	STRAUB, et al., "Genotyping Cryptosporidium parvum with an hsp 70 single-nucleotide polymorphism microarray", Applied and Environmental Microbiology, (2002) p. 1817-1826, Vol. 68.	
	C28	TAGUCHI, H., et al., "Single-molecule observation of protein-protein interactions in the chaperonin system", Nature Biotechnology, (2001) p. 861-865, Vol. 19.	

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Signature	Considered	

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Examiner Initials *	Cite No. 1		
	C29	TELL, et al. "Comparison of four rapid DNA extraction techniques for conventional polymerase chain reaction testing of three mycobacterium spp. that affects birds", Avian Diseases, (Oct. 2003) p. 1486-1490, Vol. 47.	
	C30	TIJSSEN, "Overview of principles of hybridization and the strategy of nucleic acid probe assays", Laboratory Techniques in Biochemistry and Molecular Biology Hybridization with Nucleic Acid Probes, (1993) Part I, Chapter 2, Elsevier, New York.	
	C31	WANG, Z., et al., "Basis of Substrate Binding by the Chaperonin GroEL", Biochem., (1999) p. 12537-12546, Vol. 38.	
	C32	XU, Z., et al., "The crystal structure of the asymmetric GroEL-GroES-(ADP), chaperonin complex", Nature, (1997) p. 741-750, Vol. 388.	
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